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RAW SEQUENCE LISTING

DATE: 03/13/2002

PATENT APPLICATION: US/09/993,292A

TIME: 10:59:50

Input Set : A:\UOFMD.007A.SEQLIST.TXT

Output Set: N:\CRF3\03132002\I993292A.raw

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4 <110> APPLICANT: James E. Galen
5 University of Maryland
7 <120> TITLE OF INVENTION: USE OF CLY A HEMOLYSIN FOR EXCRETION OF
8 PROTEINS
10 <130> FILE REFERENCE: UOFMD.007A
12 <140> CURRENT APPLICATION NUMBER: 09/993,292A
13 <141> CURRENT FILING DATE: 2001-11-23
15 <150> PRIOR APPLICATION NUMBER: 60/252,516
16 <151> PRIOR FILING DATE: 2000-11-22
18 <160> NUMBER OF SEQ ID NOS: 19
20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 6271
24 <212> TYPE: DNA
25 <213> ORGANISM: Artificial Sequence
27 <220> FEATURE:
28 <223> OTHER INFORMATION: pSEC84 Expression Plasmid
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33 taaagttaat gatgatagcg ggagttattc tagttgagag tgaaggtttt gttttgacat 180
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42 gttttaaaaca ggagtactcg caggaagctt ctgttttagt tggatgatt aaagttttgc 720
43 ttatggacag ccaggacaag tttttgaaag cgacacaaac tgtttatgaa tgggtgtggtg 780
44 tcgtgaogca attactctca gcgtatattt tactatttga tgaatataat gagaaaaaag 840
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133 ctggataaag ggaactcaat caagttattt tcttaccagt cattacataa tcgttattat 6180
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137 <210> SEQ ID NO: 2

138 <211> LENGTH: 305

139 <212> TYPE: PRT

140 <213> ORGANISM: Salmonella Typhi

142 <400> SEQUENCE: 2

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144 1 5 10 15
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146 20 25 30
147 Gln Val Ile Pro Trp Lys Thr Phe Asp Glu Thr Ile Lys Glu Leu Ser
148 35 40 45
149 Arg Phe Lys Gln Glu Tyr Ser Gln Glu Ala Ser Val Leu Val Gly Asp
150 50 55 60
151 Ile Lys Val Leu Leu Met Asp Ser Gln Asp Lys Tyr Phe Glu Ala Thr
152 65 70 75 80
153 Gln Thr Val Tyr Glu Trp Cys Gly Val Val Thr Gln Leu Leu Ser Ala
154 85 90 95
155 Tyr Ile Leu Leu Phe Asp Glu Tyr Asn Glu Lys Lys Ala Ser Ala Gln

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156          100          105          110
157 Lys Asp Ile Leu Ile Arg Ile Leu Asp Asp Gly Val Lys Lys Leu Asn
158          115          120          125
159 Glu Ala Gln Lys Ser Leu Leu Thr Ser Ser Gln Ser Phe Asn Asn Ala
160          130          135          140
161 Ser Gly Lys Leu Leu Ala Leu Asp Ser Gln Leu Thr Asn Asp Phe Ser
162 145          150          155          160
163 Glu Lys Ser Ser Tyr Phe Gln Ser Gln Val Asp Arg Ile Arg Lys Glu
164          165          170          175
165 Ala Tyr Ala Gly Ala Ala Ala Gly Ile Val Ala Gly Pro Phe Gly Leu
166          180          185          190
167 Ile Ile Ser Tyr Ser Ile Ala Ala Gly Val Ile Glu Gly Lys Leu Ile
168          195          200          205
169 Pro Glu Leu Asn Asn Arg Leu Lys Thr Val Gln Asn Phe Phe Thr Ser
170          210          215          220
171 Leu Ser Ala Thr Val Lys Gln Ala Asn Lys Asp Ile Asp Ala Ala Lys
172 225          230          235          240
173 Leu Lys Leu Ala Thr Glu Ile Ala Ala Ile Gly Glu Ile Lys Thr Glu
174          245          250          255
175 Thr Glu Thr Thr Arg Phe Tyr Val Asp Tyr Asp Asp Leu Met Leu Ser
176          260          265          270
177 Leu Leu Lys Gly Ala Ala Lys Lys Met Ile Asn Thr Cys Asn Glu Tyr
178          275          280          285
179 Gln Gln Arg His Gly Lys Lys Thr Leu Phe Glu Val Pro Asp Val Ala
180          290          295          300
181 Ser
182 305
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186 <211> LENGTH: 102
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223> OTHER INFORMATION: Cloning primer
193 <400> SEQUENCE: 3
194 ggatccaaaa taaggaggaa aaaaaaatga ctagtatttt tgcagaacaa actgtagagg 60
195 tagttaaaaag cgcgatcgaa accgcagatg gggcattaga tc 102
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198 <211> LENGTH: 101
199 <212> TYPE: DNA
200 <213> ORGANISM: Artificial Sequence
202 <220> FEATURE:
203 <223> OTHER INFORMATION: Cloning primer
205 <400> SEQUENCE: 4
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207 ttggtattca ttacaggtgt taatcatttt ctttcagct c 101
209 <210> SEQ ID NO: 5
210 <211> LENGTH: 97
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence

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214 <220> FEATURE:
215 <223> OTHER INFORMATION: Cloning primer
217 <400> SEQUENCE: 5
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221 <210> SEQ ID NO: 6
222 <211> LENGTH: 69
223 <212> TYPE: DNA
224 <213> ORGANISM: Artificial Sequence
226 <220> FEATURE:
227 <223> OTHER INFORMATION: Cloning primer
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231 cgcaacgcg 69
233 <210> SEQ ID NO: 7
234 <211> LENGTH: 60
235 <212> TYPE: DNA
236 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Cloning primer
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245 <210> SEQ ID NO: 8
246 <211> LENGTH: 101
247 <212> TYPE: DNA
248 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: Cloning primer
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257 <210> SEQ ID NO: 9
258 <211> LENGTH: 101
259 <212> TYPE: DNA
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263 <223> OTHER INFORMATION: Cloning primer
265 <400> SEQUENCE: 9
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267 ctagctcatg tttgacagct tatcatcgat aacctttaat g 101
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270 <211> LENGTH: 71
271 <212> TYPE: DNA
272 <213> ORGANISM: Artificial Sequence
274 <220> FEATURE:
275 <223> OTHER INFORMATION: Cloning primer
277 <400> SEQUENCE: 10
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VERIFICATION SUMMARY

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Input Set : A:\UOFMD.007A.SEQLIST.TXT

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L:260 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9

L:531 M:351 W: Sequence data Name/Key Feature Out-of-Range, SEQ ID#:19, CDS LOCATION: (0)...
(2253)